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Figure 5a

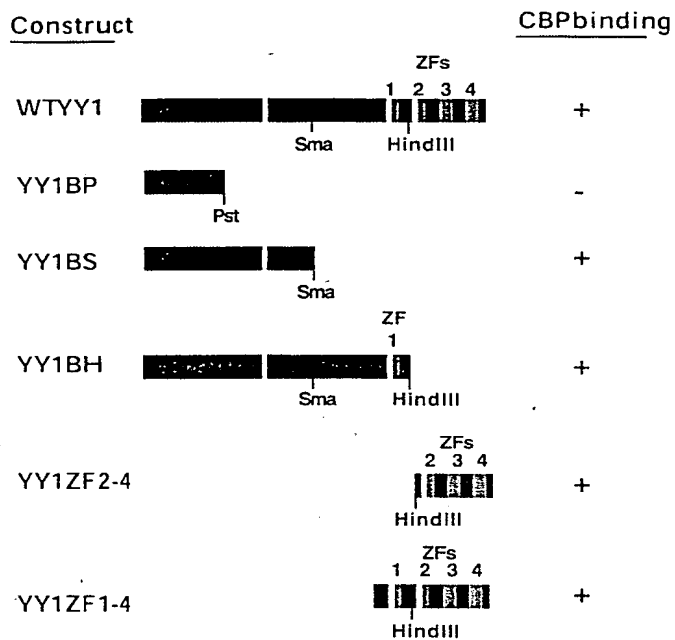


Figure 5b

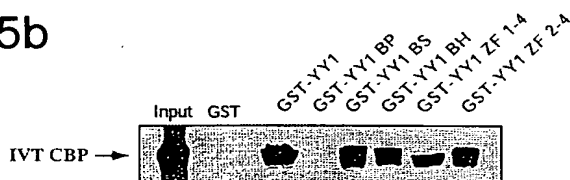
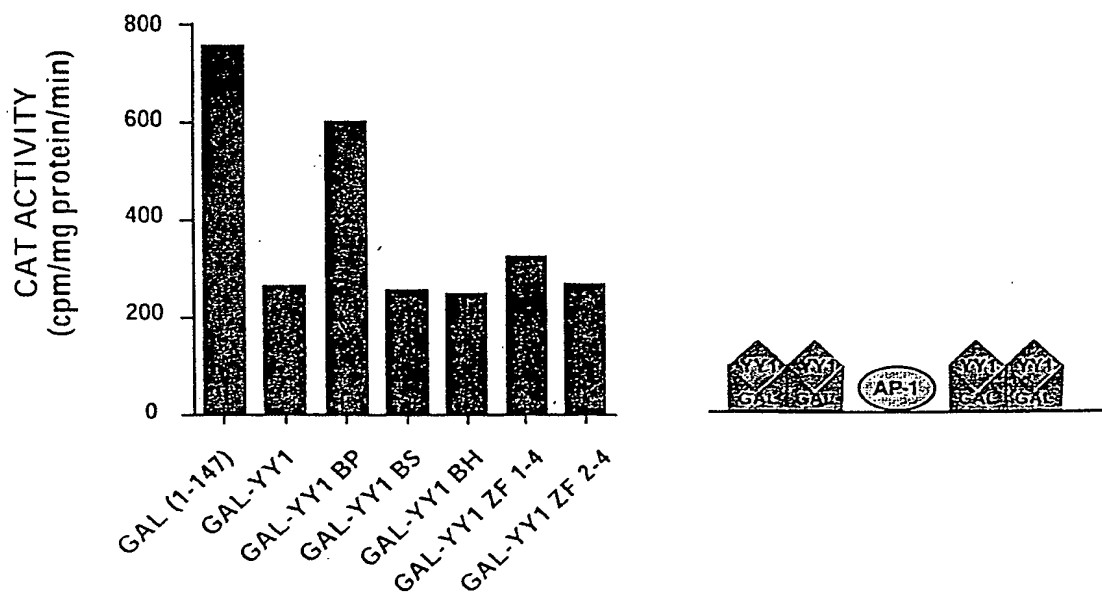
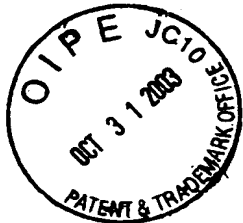


Figure 5c



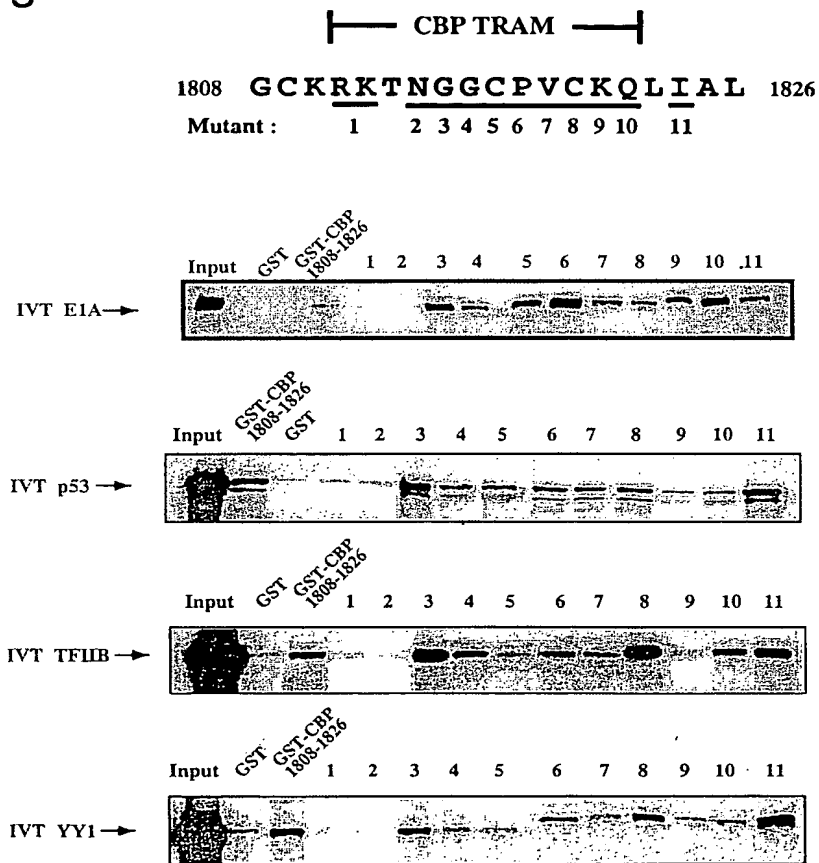


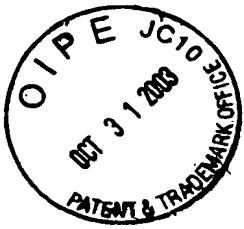
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Figure 6a

TRIMs	[	E1A	VNEFF <u>F</u> PESLILAA
		p53	SQET <u>F</u> SDLWKLLP
		E2F	FDCD <u>F</u> GDLTPLDF
		TFIIB	MMNA <u>F</u> KEITTMAD
		YY1	AEDG <u>F</u> EDQILIPV
		YY1	CTKM <u>F</u> RDNSAMRK
		YY1	CGKA <u>F</u> VESSKLKR
		MyoD	TTDD <u>F</u> YDDPCFDS

Figure 6b





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Figure 7a

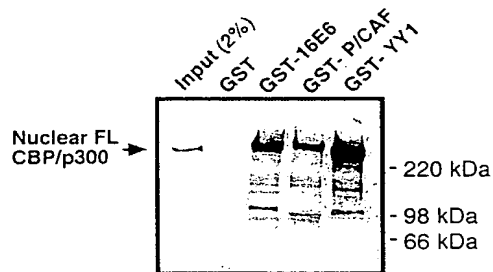


Figure 7b

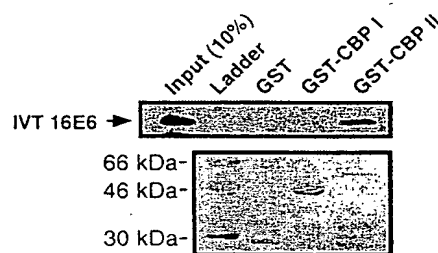


Figure 7c

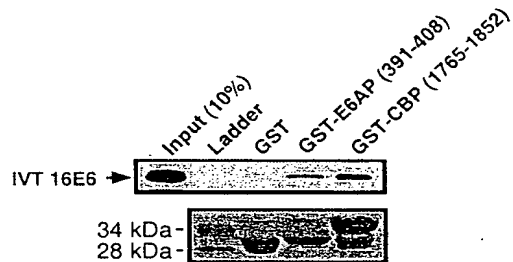
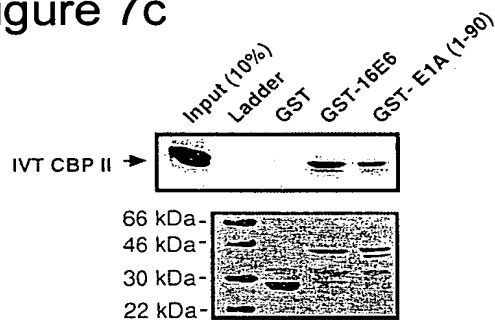
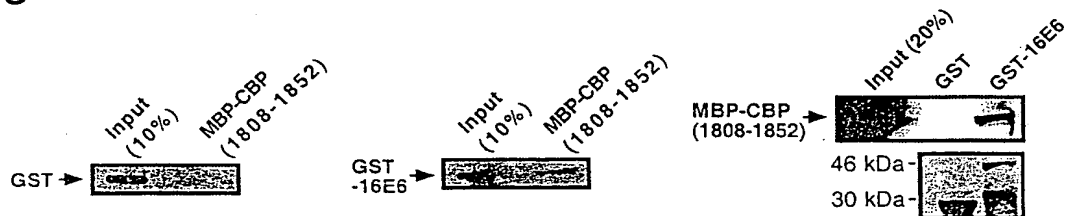
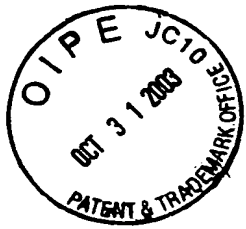


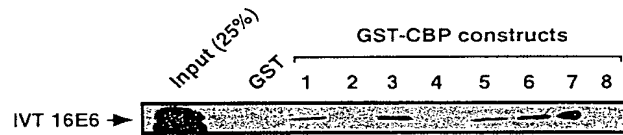
Figure 7d





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	CBP constructs	E6 binding
1	1715 1870	+
2	1715 1764	-
3	1765 1852	+
4	1765 1808	-
5	1808 1852	+
6	1808 1833	+
7	1808 1826	+
8	1808 1819	-



CBP (1808-1826): GCKRKTNGGCPVCKQLIAL  
 p300 (1770-1788): GCKRKTNGGCPICKQLIAL  
 \*\*\*\*\*+\*\*\*\*\*

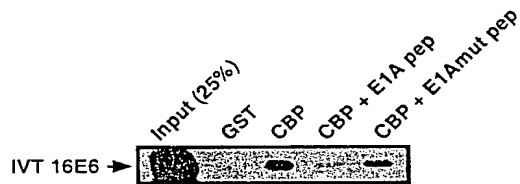
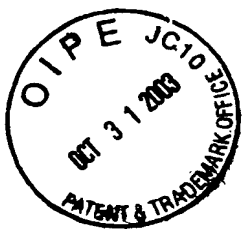


Figure 8



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Figure 9a

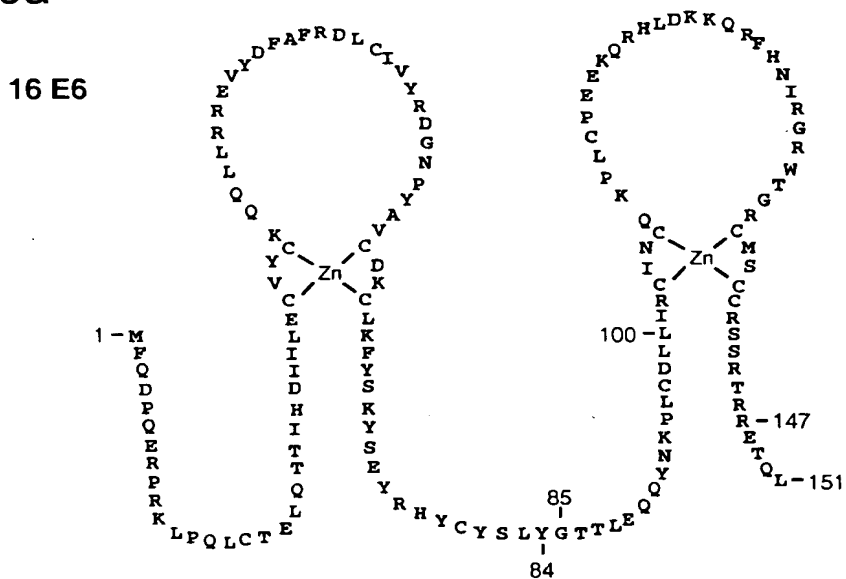


Figure 9b

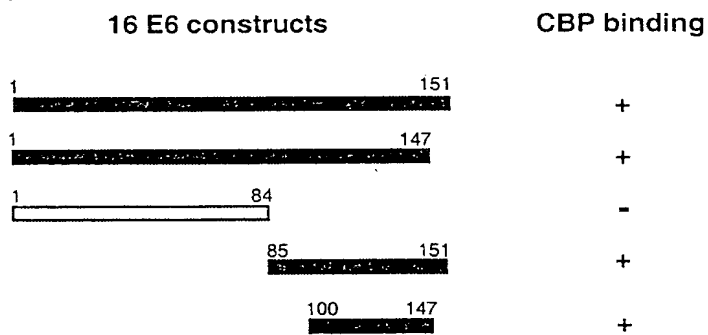
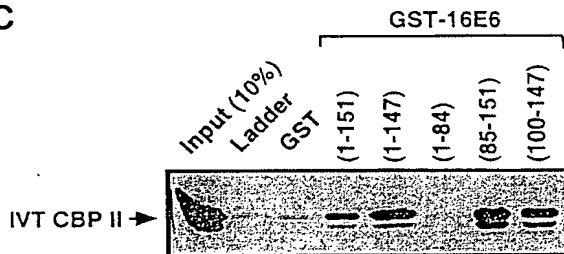
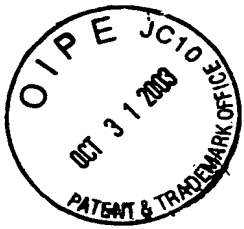


Figure 9c





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Figure 10a

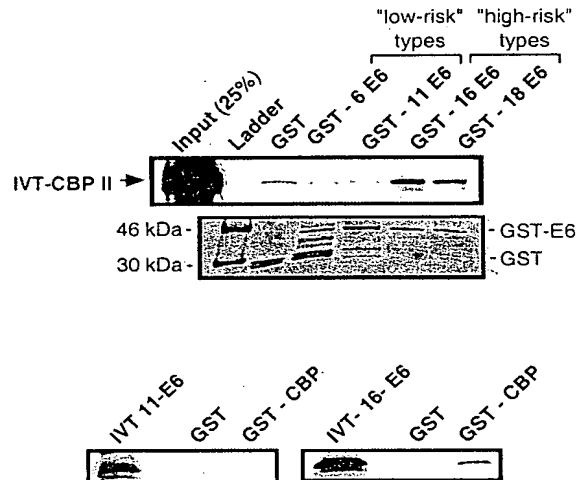


Figure 10b

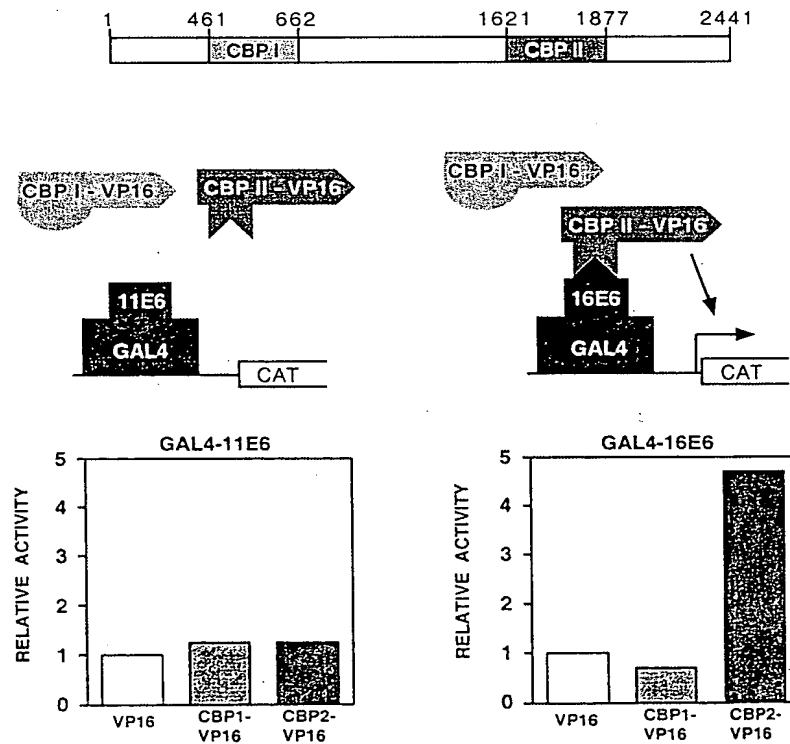




Figure 11a

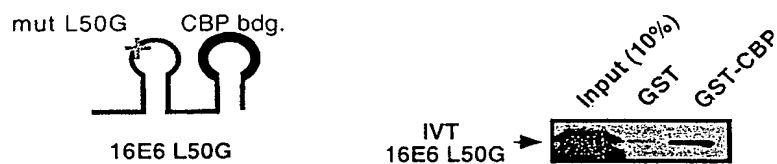


Figure 11b

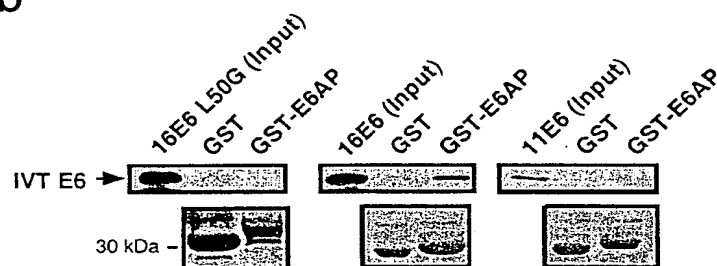
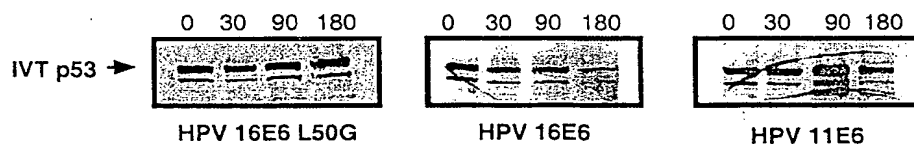
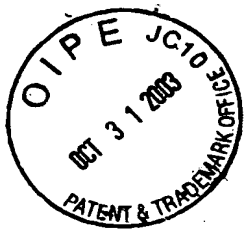


Figure 11c





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Figure 12a

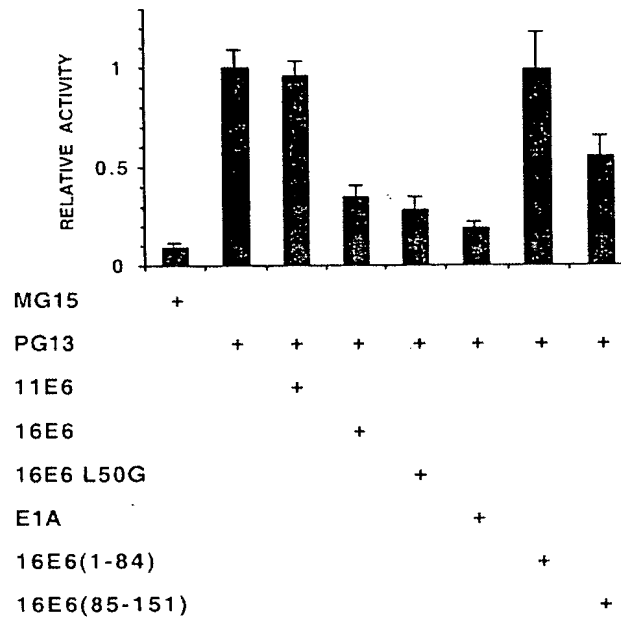
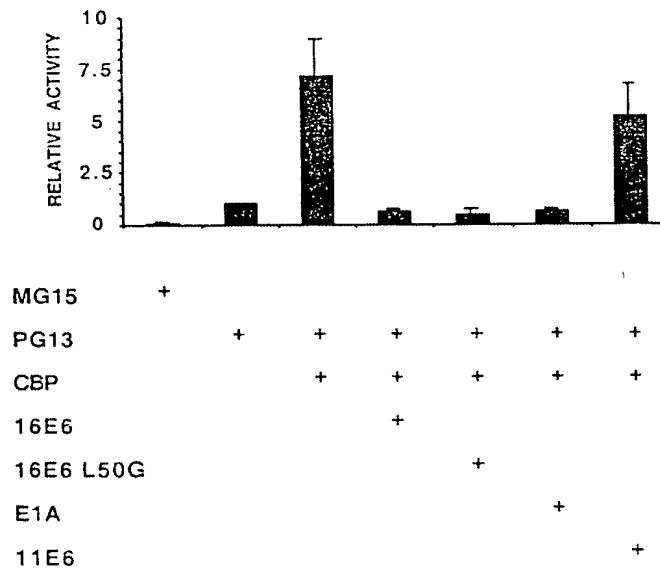
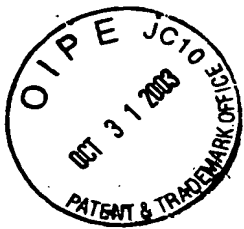


Figure 12b







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G S V D L E P G V Q K L x x x  
gGA TCC GTC GAC CTC GAG CCC GGG CTG CAG AAG CTT GAT TGA Tta gct t 3'  
cct agG CAG CTG GAG CTC GGG CCC GAC GTC TTC GAA CTA ACT AAT GGA a 5'  
BamH1 Sal1 Xho1 Pst1 HindIII stop (x3) [core HindIII]

Figure 13